## Amendments to the Specification:

Please replace the title with the following amended title:
A PURIFIED RETROVIRAL ENVELOPE POLYPEPTIDE, ISOLATED NUCLEIC
ACIDS ENCODING SAID POLYPEPTIDE, VECTORS AND USE THEREOF

Please replace the paragraph at page 6, beginning at line 27, with the following amended paragraph:

The NCBI Basic Local Alignment Search Tool (BLAST) is available from several sources, including the National Center for Biotechnology Information (NBCI, Bethesda, Md.) And on the Internet, for use in connection with the sequence analysis programs blastp, blastn, blastx, tblastn and tblastx. It can be accessed at <a href="http://www.ncbi.nlm.nih.gov/BLAST/">http://www.ncbi.nlm.nih.gov/BLAST/</a>. A description of how to determine sequence identity using this program is available at <a href="http://www.ncbi.nlm.nih.gov/BLAST/blast-help.html">http://www.ncbi.nlm.nih.gov/BLAST/blast-help.html</a>.

Please replace the paragraphs at page 21, lines 9 and 11, with the following amended paragraphs:

The mouse URL:

<http://www[[.]]DOTncbi[[.]]DOTnlm[[.]]DOTnih[[.]]DOTgov/Locus
Link/LocRpt.cgi?!=19775>

The human URL:  $\frac{1}{DOT}$ ncbi[[.]] $\frac{DOT}$ nlm[[.]] $\frac{DOT}$ nih[[.]] $\frac{DOT}$ nih[[.]] $\frac{DOT}$ gov/Locus Link/LocRpt.cqi?!=9213>

(These URLs have been inactivated by replacement of "." with "DOT".)

Please amend the paragraphs at page 21, lines 13-19 by replacing them with the following amended paragraphs:

Figure 1

Alignment of amino acid sequences of selected MLVs (from top to bottom, SEQ ID NOs:22-52) in the VR3 region. Residues corresponding to arginine 212 are shown in the rectangular box.

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Figure 2

Amino acid aligning of SL3-2 <u>(SEQ ID NO:2)</u> and MCF 247 <u>(SEQ ID NO:53)</u> envelopes: The non-homologous regions of SU, including VR3 region, are marked by brackets.

Please amend the paragraph at page 24, lines 21-22 by replacing it with the following amended paragraph:

The primer (111019: AACAATTTCACACAGGAAACAGC, SEO ID NO:3) was used as the end primer in all reactions. Other primers used are shown in table 1.

Please amend the paragraphs at page 25, line 1 to page 26, line 6 (end of table 2) by replacing them with the following amended paragraphs:

The primer (111019 :AACAATTTCACACAGGAAACAGC <u>SEQ ID NO:3</u>) was used as the end primer in all reactions. Other primers used are shown in table 1.

Table 1: Downstream primers

Library	Drimer name			
птртату	TITUEL HANG	Primer Sequence		
GI 2 0 (MGT	D0140D01	CMMMACACACACACACACACACACACACACACACACACA		
SL3-2/MCF	B8142D01	GTTTGCCCCGGTCACACTGTGCCAACAGGGTGT		
VRA	i	GGAGGCCGAGA <u>SEO ID NO:4</u>		
SL3-2/MCF	A0341A02	CTTAAGCGAGGAAACACTCCTCAGAATCAGGGC		
VRB		CCCTGTTATGATTCCTCAGCGGTCTCCAGTGAC		
		ATCAAGGGTGCCACACCGGGGGGTCGA <u>SEQ</u>		
		ID NO:5		
SL3-2/MCF	A0341A01	GGTAAAAGGGCCAGCTGGGACGGCCCCAAAGTA		
VR3		TGGGGACTAAGACTGTACCGATCCACAGGGATC		
		GACCCGGTGACCCGGTTCTCT SEQ ID NO:6		
SL3-2/MCF	B8142D02	CTAATAGTCCTGGGAATCTTAATAAGGGCAGGA		
Leader		GTATCAGTACAACATGACAGC SEQ ID NO:7		
SL3-2/MCF	A0571A05	GGTAAAAGGGCCAGCTGGGACGSNYCNAAAGYN		
alternatin		TGGGGACTAAGACTGTACCGATCCACARGRAYH		
g VR3		GACCCGGTGACCCGGTTCTCT SEO ID NO:8		
library		Control Street, and Contro		
SL3-2/MCF	T3130F07	GGTAAAAGGGCCAGCTGGGACGCCTCCAAAGCA		
VR3 RT		TGGGGACTAAGACTGTACCGATCCACAAGGACC		
		GACCCGGTGACCCGGTTCTCT SEO ID NO:9		
SL3-2/MCF	T3130F08	GGTAAAAGGGCCAGCTGGGACGCCTCCAAAGCA		
VR3 RI		TGGGGACTAAGACTGTACCGATCCACAAGGATC		
		GACCCGGTGACCCGGTTCTCT SEQ ID		
	A	NO:10		
SL3-2/MCF	T3130F09	GGTAAAAGGGCCAGCTGGGACGCCTCCAAAGCA		
VR3 GT		TGGGGACTAAGACTGTACCGATCCACAGGGACC		
		GACCCGGTGACCCGGTTCTCT SEQ ID		
		NO:11		
SL3-2/MCF	T3130F10	GGTAAAAGGGCCAGCTGGGACGCCTCCAAAGCA		
VR3 GI		TGGGGACTAAGACTGTACCGATCCACAGGGATC		
1113 01				
		GACCCGGTGACCCGGTTCTCT SEQ ID		
		<u>NO:12</u>		

## Upstream fragments

The upstream fragments were made using the primer (111020: GATTAAGTTGGGT AAGCCAGGG <u>SEO ID NO:13</u>) as the end primer. Other primers used are shown in table 2. PCR conditions were otherwise the same as those for downstream fragments.

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Table 2: Upstream primers

Construct	Primer name	
SL3-2/MCF VRA	B8087E02	CACAGTGTGACCGGGGCAAAC
		SEQ ID NO:14
SL3-2/MCF VRB	B8087E04	AGGGAGTGTTTCCTCGCTTAAG
		SEQ ID NO:15
SL3-2/MCF VR3	B8087E03	GTCCCAGCTGGCCCTTTTACC
		SEQ ID NO:16
SL3-2/MCF	B8087E05	TAAGATTCCCAGGACTATTAG
Leader		SEQ ID NO:17

Please insert the following paragraph at page 45 after line 2:

Towers et al 1999, PNAS vol. 97, no. 22 pp 12295-12299.